## SEQUENCE LISTING

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<110> Vogeli, Gabriel
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    Merchant, Kalpana
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cca ttg tgc Pro Leu Cys 35	tcc ctc Ser Leu	tac ctt Tyr Leu	att Ile 40	gct Ala	gtg Val	cta Leu	ggt Gly	aac Asn 45	ttg Leu	aca Thr	atc Ile	144
atc tac att Ile Tyr Ile 50			His									192
ttt ctt tgc Phe Leu Cys 65												240
atg ccc aaa Met Pro Lys	atg ctg Met Leu 85	gcc atc Ala Ile	ttc Phe	tgg Trp	ttc Phe 90	aat Asn	tcc Ser	act Thr	acc Thr	atc Ile 95	Gln	288
ttt gat gct Phe Asp Ala												336
atg gaa tcc Met Glu Ser 115												384
atc tgt cac Ile Cys His 130	cca ctg Pro Leu	cgc cat Arg His 135	Ala	aca Thr	gta Val	ctt Leu	acg Thr 140	ttg Leu	cct Pro	cgt Arg	gtc Val	432
acc aaa att Thr Lys Ile 145												480
ccc ctt cct Pro Leu Pro												528

					tgc Cys									Ala		576
					aat Asn											624
Āla					tca Ser											672
	Lys				ggc Gly 230											720
					cat His											768
					atg Met											816
					ttg Leu											864
 ctc. Leu	aac Asn 290	cca Pro	att. Ile	gtc Val	tat Tyr	gga Gly 295	gtg Val	aag Lys	aca Thr	aag Lys	gag Glu 300	att Ile	cga, Arg	cag Gln	cgc Arg	912
atc Ile 305	ctt Leu	cga Arg	ctt Leu	ttc Phe	cat His 310	gtg Val	gcc Ala	aca Thr	cac His	gct Ala 315	tca Ser	gag Glu	ccc Pro	tag		957
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Leu	Ile	Gly	Leu 20		Gly	Leu	Glu	Glu 25	Ala	Gln	Phe	Trp	Leu 30	Ala	Phe	
Pro		Cys 35	Ser	Leu	Tyr	Leu	Ile 40	Ala	Val	Leu	Gly	Asn 45	Leu	Thr	Ile	
Ile	Tyr 50	Ile	Val	Arg	Thr	Glu 55	Hiŝ	Ser	Leu	His	Glu 60	Pro	Met	Tyr	Ile	
Phe 65	Leu	Cys	Met	Leu	Ser 70	Gly	·Ile	Asp	Ile	Leu 75	Ile	Ser	Thr	Ser	Ser 80	
Met	Pro	Lys	Met.	Leu 85	Ala	Ile	Phe	Trp	Phe 90	Asn	Ser	Thr	Thr	.Ile 95	Gln	
Phe	Ąsp	Ala	Cys 100	Leu	Leu	Gln	Met	Phe 105	Ala	Ile	His	Ser	Leu 110	Ser	Gly	

Met	Glu	Ser 115	Thr	Val	Leu	Leu	Ala 120	Met	Ala	Phe	Asp	Arg 125	Tyr	Val	Ala	
Ile	Cys 130	His	Pro	Leu	Arg	His 135	Ala	Thr	Val	Leu	Thr 140	Leu	Pro	Arg	Val	•
Thr 145	Lys	Ile	Gly	Val	Ala 150	Ala	Val	Val	Arg	Gly 155		Ala	Leu	Met	Ala 160	
Pro	Leu	Pro	Val	Phe 165	Ile	Lys	Gln	Leu	Pro 170	Phe	Cys	Arg	Ser	Asn 175	Ile	•
Leu	Ser	His	Ser 180	Tyr	Cys	Leu	His	Gln 185	Asp	Val	Met	Lys	Leu 190	Ala	Cys	
Asp	Asp	Ile 195	Arg	Val	Asn	Val	Val 200	Tyr	Gly	Leu	Ile	Val 205	Ile	Ile	Ser	
Ala	Ile 210	Gly	Leu	Asp	Ser	Leu 215	Leu	Ile	Ser	Phe	Ser 220	Tyr	Leu	Leu	Ile	•
Leu 225	Lys	Thr	Val	Leu	Gly 230		Thr	Arg	Glu	Ala 235	Gln	Ala	Lys	Ala	Phe 240	* *
Gly	Thr	Cys	Val	Ser 245	His	Val	Cys	Ala	Val 250	Phe	Ile	Phe	Tyr	Val 255	Pro	
 Phe	Ile	Gly	Leu 260	Ser	Met	Val		Arg 265	Phe	Ser	Lys	Arg	Arg 270	Asp	Ser	
Pro	Leu	Pro 275	Val	Ile	Leu	Ala	Asn 280	Ile	Tyr	Leu	Leu	Val 285	Pro	Pro	Val	See
Leu	Asn 290	Pro	Ile	Val		Gly 295	Val	Lys	Thr	Lys	Glu 300	Ile	Arg	Gln	Arg	
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atg		agc						ata Ile								48
								ctc Leu 25						Val		96
								gga Gly								144
								gcc Ala								192

									tac Tyr								240
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									cac His 105							gga Gly	336
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									ttt Phe								480
	_								tgt Cys				_	-	_		528
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									ttc Phe								624
			Phe						tcc Ser								672
									aaa Lys								720
-									ttc Phe								768
			Thr	_	Pro				ttc Phe 265		_	_	_		_		816
									ttg Leu								864
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Leu Phe His Thr Val Ile Phe Pro Leu Leu Asn Pro Val Ile Tyr Thr

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999 90	act Thr	ctg Leu	act Thr	tgc Cys	aaa Lys 95	gtg Val	att Ile	gcc Ala	ttt Phe	ctg Leu 100	G1y	gťt Val	ttg Leu	tcc Ser	tgt Cys 105	580
								tgc Cys							Leu	628
								aca Thr 130								676
	Leu							tgg Trp							gca Ala	724
					Asp			act Thr								772
								tcc Ser								820
Gly	Phe	Met	Leu	Leu 190	Leu	Ala	Leu	Ile	Leu 195	Leu	Āla	Thr	Gln	Leu 200		868
tac Tyr	ctc Leu	Lys	ctg Leu 205	ata Ile	ttt Phe	Phe	gtc Val	cac His 210	gat Asp	cga Arg	aga Arg	aaa Lys	atg Met 215	aag Lys	cca Pro	916
								cag Gln								964
	Āla			cag			qcc	aat	taa		aca	orda.	+++	das	200	1012
	235	Ser	Gly	Gln	Ala	Ala 240	Ala	Asn								
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 Gly 250 aca	ccc Pro	aca Thr	cca Pro	ccc Pro	acc Thr 255	240 ttg Leu ttg	Ala ctg Leu	Asn	Trp atc Ile	agg Arg 260	Ala 245 caa Gln	Gly aat Asn	Phe gca Ala atg	Gly aac Asn	acc Thr 265	1060
 Gly 250 aca Thr	ccc Pro	aca Thr aga Arg	cca Pro aga Arg	ccc Pro agg Arg 270	acc Thr 255 cta Leu	ttg Leu ttg Leu	Ala ctg Leu gtc Val	Asn ggc Gly tta	Trp atc Ile gac Asp 275 act	agg Arg 260 gag Glu	Ala 245 caa Gln ttc Phe	Gly aat Asn aaa Lys	Phe gca Ala atg Met	Gly aac Asn gag Glu 280 acc	acc Thr 265 aaa Lys	
Gly 250 aca Thr aga Arg	ccc Pro ggc Gly atc Ile	aca Thr aga Arg agc Ser	cca Pro aga Arg aga Arg 285	agg Arg 270 atg Met	acc Thr 255 cta Leu ttc Phe	240 ttg Leu ttg Leu tat Tyr	Ala ctg Leu gtc Val ata Ile	Asn ggc Gly tta Leu atg Met 290 tat Tyr	atc Ile gac Asp 275 act Thr	agg Arg 260 gag Glu ttt Phe	Ala 245 caa Gln ttc Phe ctg Leu	Gly aat Asn aaa Lys ttt Phe	gca Ala atg Met cta Leu 295	aac Asn gag Glu 280 acc Thr	acc Thr 265 aaa Lys ttg Leu	1108

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7		Gln									att Ile 340						1300
											tac Tyr						1348
				_			_	gtt Val	ata Ile 370	tgag	39			, ,	•		1380
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		•		20		,	* T	* - , *	25		Leu			30	٠.		
-			35			٠.		40			Ile Leu	. : '	45	•	•		
		50					55				Pro 75	60					
1		Lys	Asn	Gly	Ser 85	Thr	Trp	Thr	Tyr	Gly 90	Thr	Leu	Thr	Cys	Lys 95	- 1	
		,		100	•				105		His			110			
;	4		115	r ii e				120			Ile		125				
7		130 Trp	. '				135 Ala				Pro 155	140	_	٠.			
	:		Tyr	Ser	Phe 165			Glu	Glu	Asp 170	Gln	Cys	Thr	Phe	Gln 175		
				180	• .		•		185		Phe			190			. *
			195				٠.	200		<del>-</del>	Leu		205				
S		2.10	-			-	215				Ala 235	220					
•					-	230					233					~ · ·	

Ala Asn Trp Leu Ala Gly Phe Gly Arg Gly Pro Thr Pro Pro Thr Leu 245 250 Leu Gly Ile Arg Gln Asn Ala Asn Thr Thr Gly Arg Arg Leu Leu Val Leu Asp Glu Phe Lys Met Glu Lys Arg Ile Ser Arg Met Phe Tyr Ile Met Thr Phe Leu Phe Leu Thr Leu Trp Gly Pro Tyr Leu Val Ala ,295 Cys Tyr Trp Arg Val Phe Ala Arg Gly Pro Val Val Pro Gly Gly Phe 310 315 Leu Thr Ala Ala Val Trp Met Ser Phe Ala Gln Ala Gly Ile Asn Pro Phe Val Cys Ile Phe Ser Asn Arg Glu Leu Arg Arg Cys Phe Ser Thr 345 Thr Leu Leu Tyr Cys Arg Lys Ser Arg Leu Pro Arg Glu Pro Tyr Cys 365 Val Ile ..370 <210> 15 <211> 1191 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1) .. (1188) atg ttt aga cct ctt gtg aat ctc tct cac ata tat ttt aag aaa ttc Met Phe Arg Pro Leu Val Asn Leu Ser His Ile Tyr Phe Lys Lys Phe 10 cag tac tgt ggg tat gca cca cat gtt cgc agc tgt aaa cca aac act 96 Gln Tyr Cys Gly Tyr Ala Pro His Val Arg Ser Cys Lys Pro Asn Thr gat gga att tca tct cta gag aat ctc ttg gca agc att att cag aga Asp Gly Ile Ser Ser Leu Glu Asn Leu Leu Ala Ser Ile Ile Gln Arg 3.5 gta ttt gtc tgg gtt gta tct gca gtt acc tgc ttt gga aac att ttt 192 Val Phe Val Trp Val Val Ser Ala Val Thr Cys Phe Gly Asn Ile Phe .50 55 gtc att tgc atg cga cct tat atc agg tct gag aac aag ctg tat gcc Val Ile Cys Met Arg Pro Tyr Ile Arg Ser Glu Asn Lys Leu Tyr Ala 70 65 atq tea atc att tet etc tqc tqt qec qae tqc tta atq qqa ata tat Met Ser Ile Ile Ser Leu Cys Cys Ala Asp Cys Leu Met Gly Ile Tyr 90

,			100													
t t Le	a ttc u Phe	gtg Val	atc Ile 100	gga Gly	ggc Gly	ttt Phe	gac Asp	cta Leu 105	aag Lys	ttt Phe	cgt Arg	gga Gly	gaa Glu 110	tac Tyr	aat Asn	336
aa Ly	g cat s His	gcg Ala 115	cag Gln	ctg Leu	tgg Trp	atg Met	gag Glu 120	agt Ser	act Thr	cat His	tgt Cys	cag Gln 125	ctt Leu	gta Val	gga Gly	384
	t ttg r Leu 130	Ala														432
ct Le 14	g aca u Thr 5	ttg Leu	gaa Glu	aaa Lys	tac Tyr 150	atc Ile	tgc Cys	att Ile	gtc Val	tat Tyr 155	cct Pro	ttt Phe	aga Arg	tgt Cys	gtg Val 160	480
	a cct g Pro															528
	t ggt r Gly	Phe														576
	a aac s Asn															624
	t aca p Thr 210	Glu														672
	t att y Ile 5															720
	g ttt t Phe															768
	t caa n Gln															816
	a ttt l Phe							·Ile								864
ct Le	t tca u Ser 290	Leu	ctt Leu	cag Gln	gta Val	gaa Glu 295	ata Ile	cca Pro	ggt Gly	acc Thr	ata Ile 300	acc Thr	tct Ser	tgg Trp	gta Val	912
Va	g att l Ile 5			_				_	_	_						960
	t ctg r Leu			_				<del></del> -	-							1008
	c tac n Tyr															1056

									1										
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									ttc Phe									1152	
		Ile							aat Asn				tga			•		1191	
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		<i>'</i> .		·															
		)> 16 Phe		Pro	Leu 5	Val	Asn	Leu	Ser	His 10	Ile	Tyr	Phe	Lys	Lys 15	Phe			
	Gln	Tyr	Cys	Gly 20	Tyr	Ala	Pro	His	Val 25	Arg	Ser	Cys	Lys	Pro 30	Asn	Thr			
	Asp.	Gly	Ile 35	Ser	Ser	Leu	Glu	Asn 40	Leu	Leu	Ala	Ser	Ile. 45	Ile	Glņ	Arg	•		
	Val	Phe 50	Val	Trp	Val	Val	Ser 55	Ala	Val	Thr	Cys	Phe 60	Gly	Asn	Ile	Phe			
;	Val 65	Ile	Cys	Met	Arg	Pro 70	Tyr	Ile	Arg	Ser	Glu 75	Asn	Lys	Leu	Tyr	Ala 80			
	Met	Ser	Ile	Ile	Ser 85	Leu	Cys	Cys	Ala.	Asp 90	Cys	Leu	Met	Gly	Ile 95	Tyr .	÷		
	Leu	Phe	Val	Ile 100	Gly	Gly	Phe	Asp	Leu 105	Lys	Phe	Arg	Gly	Glu 110	Tyr	Asn		<i>*</i> .	•
			115	. ; :	•			120	Ser				125						
		130					135		Val	=		140							
	145	-				150			Ile		155	•				160	•		
					165				Thr	170					175		:		
		-		180					Pro 185	;			 	190					
	-		195		_			200	Vál				205						
	_	210					215		Ile			220							
	Gly 225		Asn	Leu	Ala.	Ala 230	Phe	тте	Ile	тте	Val 235	Phe	ser	ıyr	GTÀ	Ser 240	,		

Met Phe Tyr Ser Val His Gln Ser Ala Ile Thr Ala Thr Glu Ile Arg 250 Asn Gln Val Lys Lys Glu Met Ile Leu Ala Lys Arg Phe Phe Phe Ile 265 Val Phe Thr Asp Ala Leu Cys Trp Ile Pro Ile Phe Val Val Lys Phe Leu Ser Leu Leu Gln Val Glu Ile Pro Gly Thr Ile Thr Ser Trp Val 295 Val Ile Phe île Leu Pro Ile Asn Ser Ala Leu Asn Pro Ile Leu Tyr 310 315 Thr Leu Thr Thr Arg Pro Phe Lys Glu Met Ile His Arg Phe Trp Tyr Asn Tyr Arg Gln Arg Lys Ser Met Asp Ser Lys Gly Gln Lys Thr Tyr 345 Ala Pro Ser Phe Ile Trp Val Glu Met Trp Pro Leu Gln Glu Met Pro 360 355 365 Pro Glu Leu Met Lys Pro Asp Leu Phe Thr Tyr Pro Cys Glu Met Ser 375 Leu Ile Ser Gln Ser Thr Arg Leu Asn Ser Tyr Ser 390 <210> 17 <211> 1164 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (13)..(1089) cacaactgaa ga atg ggg ttc aac ttg acg ctt gca aaa tta cca aat aac 51 Met Gly Phe Asn Leu Thr Leu Ala Lys Leu Pro Asn Asn gag ctg cac ggc caa gag agt cac aat toa ggc aac agg agc gac ggg Glu Leu His Gly Gln Glu Ser His Asn Ser Gly Asn Arg Ser Asp Gly 15 20 25 cca gga aag aac acc ctt cac aat gaa ttt gac aca att gtc ttg Pro Gly Lys Asn Thr Thr Leu His Asn Glu Phe Asp Thr Ile Val Leu cca gtg ctt tat ctc att ata ttt gtg gca agc atc ttg ctg aat ggt 195 Pro Val Leu Tyr Leu Ile Ile Phe Val Ala Ser Ile Leu Leu Asn Gly tta gca gtg tgg atc ttc ttc cac att agg aat aaa acc agc ttc ata 243 Leu Ala Val Trp Ile Phe Phe His Ile Arg Asn Lys Thr Ser Phe Ile tto tat oto aaa aac ata gtg gtt goa gao oto ata atg acg otg aca 291 Phe Tyr Leu Lys Asn Ile Val Val Ala Asp Leu Ile Met Thr Leu Thr

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	_				_	_		act Thr		_	_			_		_	387
								Gly 999									435
								gac Asp									483
								gtt Val 165									531
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								agt Ser									627
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	_			_			_	ata Ile								_	723
								agc Ser 245									771
٠								ttt Phe									819
		. —	-					ttt Phe	_			_			Leu	_	867
								tat Tyr									915
								ctg Leu									963
								ctg Leu 325									1011
•		_	_	_		_		ctg Leu		. –		_	_	_	_	_	1059

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				tat Tyr						tag	gcct	ttta	att <u>c</u>	gtttg	gttgg	ga	1109
	atico	gatai	tat a	acaaa	atat	a aa	ataaa	atatt	t tict	ettto	catt	àaaa	aaaaa	aaa a	aaaaa	a .	1164
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		0> 18 Glv		Asn	Lèu	Thr	Leu	Δla	Lvs	Len	Pro	Asn	Asn	Glu	Leu	His	
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,	Gly	Gln	Glu	Ser 20	His	Asn	Ser	Gly	Asn 25	Arg	Ser	Asp	Gly	Pro 30	GIA	Lys	
	Asn	Thr	Thr	Leu	His	Asn	Glu	Phe	Asp	Thr	Ile	Val	Leu	Pro	Val	Leu	
			35				,	40		÷			45				
	m	T 011	T 1 0	Tlo	Dho	17.7	<b>7</b> . Τ ~ .	Cox	T ] 0	T 011	T 011	Λαπ	Cirr	T 011	ח ד ת		
	тут	50	116	Ile	PHE	vai	55	ser	116	ьец	ьеи	60	GIY	Бец	Hia	vai,	•
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		Ile	Phe	Phe	His		Arg	Asn	Lys	Thr		Phe	Ile	Phe	Tyr	Leu	
	65	•	٠.,			70					75			-		80	
	Lys	Asn	Ile	Val	Val 85	Ala	Asp	Leu	Ile	Met 90	Thr	Leu	Thr	Phe	Pro 95	Phe	
	7	T 1'-		77.5		<b>3</b> 3 -		Dl				m	D1	<b>T</b>	i.	T1.	
	Arg	ше	vaı	His 100	Asp	Ата	GIY	Pņe	105	Pro	Trp	Tyr	Pne	ьуs 110	Pne	TTE	
				100					10,5	:				,			•
	Leu	Cys	Arg	Tyr	Thr	Ser	Val		Phe	Tyr	Ala	Asn	Met	Tyr	Thr	Ser	
		•	115			•		120					125				
	Ile	Val	Phe	Leu-	Glv	Leu	Ile	Ser	Ile	Asp	Ara	Tvr	Leu	Lvs	Val	Val	
		130	7.77		2		135				· .	140					
	_			~ 3	_	_	_		_	_			-	1			
	Lуs 145	Pro	Pne	Gly	Asp	5er 150	Arg	Met	Tyr	ser	11e	Thr	Pne	Thr	гÀг	vai 160	
	113		. 199			130		٠.				1.				100	
٠,	Leu	Ser	Vā İ	Cys	Val			Ile	Met	Ala	Val	Leu	Ser	Leu	Pro	Asn	
	`				165	*- 1				170					175		
	Ile	Île	Leu	Thr	Asn	Gly	Gln	Pro	Thr	Glu	Asp	Asn	Ile	His	Asp	Cys	
	٠.,			180					185		-			190	-	_	
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	Tyr		Asn	Ser	Cys	Leu		Val	Ala		Leu			Leu	Ile	Gly	
		210					215					220					
	Cys	Tyr	Ile	Ala	Ile	Ser	Arg	Tyr	Ile	His	Lys	Ser	Ser	Arg	Gln	Phe	
	225					230		•			235	,				240	
	Tla	Sèr	Gla	Sar	, Ser	Δ×~	Lve	Δra	Laze	ніа	Δαπ	Gla	Ser	Tla	Δ~~	U=1	
	116	DET.	G.T.11	Ser	245	ALY.	пур	Arg	пур	250	TOII	GIII	261	116	255	VAI	
	Val	Val	Ala		Phe	Phe	Thr	Cys		Leu	Pro	Tyr	His		Cys	Arg	
				260					265					270			,
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		•													•	
Ile	Pro	Phe 275	Thr	Phe	Ser	His	Leu 280	Asp	Arg	Ļeu ,	Leu	Asp 285	Glu	Ser	Ala	
Gln	Lys 290	Ile	Leu	Tyr	Tyr	Cys 295	Lys	Glu	Ile	Thr	Leu 300	Phe	Leu	Ser	Ala	
Cys 305	Asn	Val	Суѕ	Leu	Asp 310	Pro	Ile	Ile	Tyr	Phe 315	Phe	Met	Cys	Arg	Ser 320.	
Phe	Ser	Arg	Arg	Leu 325	Phe	Lys	Lys	Ser	Asn 330	Ile	Arg	Thr	Arg	Ser 335	Glu	
Ser	Ile	Arg	Ser 340	Leu '	Gln	Ser	Val	Arg 345	Arg	Ser	Glu	Val	Leu 350	Ile	Tyr	· **.
Tyr	Asp	Tyr 355	Thr	Asp	Val	•			•							
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•			*, **									1			5	
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Ser	Ser	Thr	Asn	Ser 10 ttg	Ser	Val gtc	Leu	Pro agc Ser	Cys 15 ttg	Pro gtg	Asp	Tyr	Arg gcc Ala	Pro 20 999	Thr	
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Ser cac His	ser cgc Arg	Thr ctg Leu	Asn cac His 25	Ser 10 ttg Leu cta	Ser gtg Val gcc	Val gtc Val ctc	tac Tyr	agc ser 30	Cys 15 ttg Leu ttc	Pro gtg Val	Asp ctg Leu cgc	Tyr gct Ala gcg	gcc Ala 35 ctg	Pro 20 999 Gly cgc	Thr ctc Leu gtg	
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Ser cac His	ser cgc Arg	Thr ctg Leu	Asn cac His 25	Ser 10 ttg Leu cta	Ser gtg Val gcc	Val gtc Val ctc	tac Tyr	agc ser 30	Cys 15 ttg Leu ttc	Pro gtg Val	Asp ctg Leu cgc	Tyr gct Ala gcg	gcc Ala 35 ctg	Pro 20 999 Gly cgc	Thr ctc Leu gtg	152
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Arg Ala Leu Arg Val His Ser Val Val Ser Val Tyr Met Cys Asn Leu 50 55 60

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Tyr Tyr Ala Leu His His Trp Pro Phe Pro Asp Leu Leu Cys Gln Thr 85 90 95

Thr Gly Ala Ile Phe Gln Met Asn Met Tyr Gly Ser Cys Ile Phe Leu 100 105 110

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